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OM protein nucleic search, using frame_plus_p2n model

Run on: January 16, 2003, 16:51:22 : Search time 212.829 Seconds
(without alignments)
137.557 Million cell updates/sec

Title: US-09-856-070-19

Perfect score: 65

Sequence: 1 KKEIMLRIDYEE 13

Scoring table: PLASUM52

Xgapop 10.0, Xgapext 0.5

Xgapop 10.0, Xgapext 0.5

Xgapop 6.0, Xgapext 7.0

Delop 6.0, Delext 7.0

Searched: 2185236 seqs, 113000000 residues

Total number of hits satisfying chosen parameters: 4370478

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Command line parameters:

-MODEL=frame_p2n model -DBV=xl
-Q=ccac2.1/195PTO, spool/090856070/runat 14012003_158433_1611/APP_query fasta_1 1592
-PR=N-Geneset-1.161002 -CRMI=fastaq -SUFFIX=ref -MINMATCH=0.1 -L=seq2-0
-LOCUS=1 -UNIT=bits -START=1 -END=1 -MAPLX=blom62 -TRANS=human43 edi
-LIST=45 -DOCALIGN=200 -THR_SCORE=ECT -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-MODE=LOCAL -OUTFMT=pro -NRM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
-USER=ins090856070/seqn_1_1449/seqn_14012003_158433_1611 -NPROG 6 -TOPIC 4
-NO_XLIFY -NO_MAP -LAPGENEPE -NEW_SCORES 0 -WAIT -L=NGLE-1 -LEV_TIME=0.120
-WARN_TIMEOUT=30 -THRAIDS=1 -VCAPEXT=0 -VCAPEXT=10 -VCAPEXT=5 -VCAPEXT=7
-XGAPOP=10 -VCAPEXT=0.5 -DELOP=6 -DELEXT=7

Database: N-Geneset_101002:*

1: /SID52/qcdata/qcseqseq/geneseq/emb1/NA1980.DAT:*

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24: /SID52/qcdata/qcseqseq/geneseq/emb1/NA2003.DAT:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Match	Query Length	DB ID	Description
1	65	100.0	2595	22	Human colon cancer
2	65	100.0	2595	24	Human osteoblast d
3	65	100.0	2595	24	Human lung cancer
4	65	100.0	3044	24	Human osteoblast d
5	65	100.0	3044	24	Human cDNA differe
6	65	100.0	3044	24	Cene #1721 used to
7	65	100.0	3047	24	Human ovarian tumor
8	65	100.0	4072	24	Human osteoblast d
9	65	100.0	3115	21	Human colon cancer
10	65	100.0	11445	22	Human immune/haema
11	41	63.1	1447	23	DNA encoding novel
12	41	63.1	66494	24	Human osteoblast d
13	43	61.5	224	22	Human fetal liver
14	40	61.5	224	22	Human brain expres
15	40	61.5	523	22	Human fetal liver
16	40	61.5	523	22	Human brain expres
17	40	61.5	1815	23	DNA encoding novel
18	40	61.5	2528	21	Human secreted pro
19	40	61.5	2701	22	Human cDNA sequenc
20	40	61.5	2991	24	Human prostate exp
21	40	61.5	2973	23	DNA encoding novel
22	40	61.5	4226	23	Human cervical can
23	40	61.5	4558	22	Human cDNA differe
24	40	61.5	149671	24	Human fetal liver
25	39	60.0	205	22	Human brain expres
26	39	60.0	205	22	Human bone marrow
27	39	60.0	205	22	Probe #20109 used
28	39	60.0	205	22	Human genome deriv
29	39	60.0	205	22	Human secreted pro
30	39	60.0	205	22	Human cDNA differe
31	39	60.0	452	22	Human fetal liver
32	39	60.0	452	22	Human brain expres
33	39	60.0	452	22	Human bone marrow
34	39	60.0	452	22	Probe #7056 used t
35	39	60.0	452	22	Human genome deriv
36	39	60.0	523	21	Human secreted pro
37	39	60.0	523	24	Human cDNA differe
38	39	60.0	592	22	Human low adenosin
39	39	60.0	1255	21	Human adenosine re
40	39	60.0	1400	13	Encodes a HeLa cel
41	39	60.0	1400	15	DNA encoding a giv
42	39	60.0	1867	22	Human colon cancer
43	39	60.0	1985	24	DNA encoding novel
44	39	60.0	2134	18	Human myeloid cell
45	39	60.0	2142	23	DNA encoding novel

ALIGNMENTS

RESULT 1

AAH33385

12 AAH33385 standard: LNA: 2595 BP.

XX AAH33385;

XX 03 Sep 2001 (first entry)

XX Human colon cancer antigen encoding cDNA SEQ ID NO:441.

XX Homo sapiens, colorectal carcinoma, ss.

XX Homo sapiens.

XX W250123920 A2.

XX 05-APR-2001.

PD

PF 20-SEP-2001; 2001WO 0542232.
 XX
 PP 22-SEP-2000; 2000RS-234837P
 PP 10-OCT-2000; 2000RS-234440P.
 PP 20-JUN-2001; 2001US-301928P.
 XX
 PA (CORI-) CORIAX CORP.
 XX
 PI Benson DP, Mohanath P, Lodes MJ;
 XX
 DR WPI: 2002 372001/40
 XX
 PI New tumor lung proteins and nucleic acids encoding the proteins, useful
 PT as vaccines and for treating, preventing, diagnosing or monitoring lung
 PT cancer
 XX
 PS Claim 1: Page 159-160, 189pp; English.
 XX
 CC The invention relates to an isolated polynucleotide comprising a sequence
 CC selected from 183 human DNA sequences (appearing as ABK70130-ABK70312),
 CC or their fragments, homologues, variants or complements and their encoded
 CC polypeptides. Also included are an expression vector comprising the
 CC polynucleotide operably linked to an expression control sequence; a host
 CC cell transformed or transfected with an expression vector; an isolated
 CC antibody, or its antigen-binding fragment that specifically binds to the
 CC polypeptide; a method for detecting the presence of a cancer in a
 CC patient; a fusion protein comprising at least the polypeptide; an
 CC oligonucleotide that hybridises to the polynucleotide under moderately
 CC stringent conditions; a method for stimulating and/or expanding T cells
 CC specific for a tumour protein; an isolated T cell population comprising T
 CC cells prepared from the method of above; a composition comprising a first
 CC component consisting of carriers and immunostimulants, and a second
 CC component selected from the polynucleotides, proteins, antibodies, fusion
 CC proteins, T cell populations and antigen presenting cells expressing the
 CC polypeptide; methods for stimulating an immune response or treating
 CC cancer in a patient by administering the composition and diagnostic kits
 CC comprising at least one of the oligonucleotide of, or an antibody and a
 CC detection reagent consisting of a reporter group. The polypeptides and
 CC polynucleotides are useful as vaccines for the treatment or prevention of
 CC lung cancer, and for diagnosis and monitoring of such cancer. The
 CC polynucleotide, polypeptide and antigen presenting cells can be
 CC used to stimulate or expand T cells specific for a tumorous protein.
 CC The polynucleotides may be used as probes or primers for nucleic acid
 CC hybridisation, and in the preparation of ribozyme molecules for
 CC inhibiting expression of tumour polypeptides and proteins in tumour
 CC cells. The present sequence is one of the 183 lung cancer associated
 CC polynucleotides.
 XX
 SQ Sequence 2930 BP, 793 A; 658 C; 821 G; 658 T; 0 other;

Alignment Scores:
 Pred. No.: 0.00545 Length: 2930
 Score: 65.00 Matches: 13
 Percent Similarity: 100.00% Conservative: 0
 Best Local Similarity: 100.00% Mismatches: 0
 Query Match: 100.00% Indels: 0
 DB: 24 Gaps: 0

US-09-856-070-19 (1-13) x ABK70285 (1-2930)

QY 1 LysGluGluMetLeuArgLeuGlnAspTyrGluGlu 13
 Db 1106 AAGGAGGATTATGTTGAGGTTGAGGATATGAGGAG 1144

RESULT 4
 ABQ88180
 ID ABQ88180 standard; cDNA: 3044 BP.
 XX
 AC ABQ88180;
 XX
 AC ABQ88180;
 XX

DT 18-SEP-2002 (first entry)

DE Human osteoblast differentiation related cDNA SEQ ID NO 87.

XX Human, osteoblast, stem cell differentiation; bone tissue deposition;
 KW osteoporosis; osteopathic; ss.
 XX
 OS Homo sapiens.
 XX
 PN WO200203031 A2.
 XX
 PD 27-JUN-2002.
 XX
 FE 18 DEC-2001, 2001WO-0518276.
 XX
 XX 18-DEC-2000; 2000US 255822P.
 PP 24-APR-2001; 2001US 285691P.
 XX
 PA (GENE-) GENE LOGIC INC.
 PA (PRO-) PRACTER & GAMBLE CO.
 XX
 FI Ji D, Axelrod DW, Cook JS, Jaiswal N, Einstein K, Houghton A;
 PI Mertz L;
 DR WPI: 2002-557663/59.
 XX
 CC use of genes and their expression profiles associated with osteoblast
 CC differentiation for screening modulators bone formation, for diagnosing
 CC or treating e.g. osteoporosis, or as markers for the differentiation
 CC process
 XX
 CC claim 1: SEQ ID NO 87; 78pp + Sequence Listing; English.
 XX
 CC the invention relates to genes and their expression profiles are used
 CC for:
 CC (a) screening modulators of precursor stem cell differentiation into
 CC osteoblasts, or bone tissue deposition;
 CC (b) diagnosing abnormal deposition of bone tissue, abnormal rate of
 CC osteoblast formation or osteoporosis; or
 CC (c) treating or monitoring treatment of the conditions cited in (b), or
 CC monitoring the progression of bone tissue deposition.
 CC Specific conditions include postmenopausal osteoporosis, glucocorticoid
 CC osteoporosis or male osteoporosis, osteopenia, osteodys trophy,
 CC drug-induced abnormalities in bone formation or bone loss, conditions
 CC that involve altered bone metabolism (e.g. idiopathic juvenile
 CC osteoporosis), skeletal disease linked to breast cancer, mastocytosis,
 CC Fanconi syndrome or fibrous dysplasia. The present sequence is that of an
 CC osteoblast differentiation associated cDNA marker of the invention.
 CC Note: The sequence data for this patent did not form part of the printed
 CC specification, but was obtained in electronic format directly from WIPO
 CC at ftp.wipo.int/pub/published_pct_sequences.
 XX
 SQ Sequence 3044 BP; 826 A; 687 C; 855 G; 675 T; 1 other;

Alignment Scores:
 Pred. No.: 0.0057 Length: 3044
 Score: 65.00 Matches: 13
 Percent Similarity: 100.00% Conservative: 0
 Best Local Similarity: 100.00% Mismatches: 0
 Query Match: 100.00% Indels: 0
 DB: 24 Gaps: 0

US-09-856-070-19 (1-13) x ABQ88180 (1-3044)

QY 1 LysGluGluMetLeuArgLeuGlnAspTyrGluGlu 13
 Db 1147 AAGGAGGATTATGTTGAGGTTGAGGATATGAGGAG 1185

RESULT 5
 ABK84552
 ID ABK84552 standard; cDNA: 3044 BP.
 XX
 AC ABK84552;
 XX
 DT 14-AUG-2002 (first entry)
 XX

Human cDNA differentially expressed in granulocytic cells #1123.

XX Human, ss: granulocytic cells; DNA chip; bacterial infection;
 XX viral infection; parasitic infection; protozoal infection;
 KW fungal infection; sterile inflammatory disease; psoriasis;
 KW rheumatoid arthritis; glomerulonephritis; asthma; thrombosis;
 KW cardiac reperfusion injury; renal reperfusion injury; AIDS;
 KW adult respiratory distress syndrome; inflammatory bowel disease;
 KW Crohn's disease; ulcerative colitis; periodontal disease;
 KW granulocyte activation; chronic inflammation; allergy.

XX Homo sapiens.

XX W0200228999-A2.

XX 11-APR-2002.

XX 03-OCT-2001; 2001WO 0530921.

XX 03-OCT-2000; 2000US-247149P.

XX (GENE-) GENE LOGIC INC.

XX Rezer Barclay Y, Weissman SM, Yamaga S, Vockley J.

XX W01: 2002-435324/46

XX Detecting granulocyte activation by detecting differential expression
 PT of genes associated with granulocyte activation, which serves as
 PT diagnostic markers that is useful for monitoring disease states and
 PT drug toxicity.

XX Claim 1; SEQ ID No 1123; 114pp; English.

XX The invention relates to detecting (M1) granulocyte (G) activation
 CC (GCA), by detecting the level of expression of gene(s) (Gs) identified by
 CC DNA chip analysis as given in the specification, and comparing
 CC the expression level to an expression level in an unactivated
 CC Gc, where differential expression of Gs is indicative of GCA.
 CC Also included are modulating (M2) Gc by contacting Gc with an agent
 CC that alters the expression of at least one gene in Gs; (2) screening (M3)
 CC for an agent capable of modulating GCA or an inflammation (especially
 CC chronic) in a tissue, an allergic response in a subject, exposure of a
 CC subject to a pathogen or sterile inflammatory disease using the
 CC gene expression profile; (3) detecting (M4) an inflammation (especially
 CC chronic) in a tissue, an allergic response in a subject, exposure of a
 CC subject to a pathogen or sterile inflammatory disease, by detecting the
 CC level of expression in a sample of the tissue of gene(s) from Gs, where
 CC the level of expression of the gene is indicative of inflammation;
 CC (4) treating (M5) an inflammation (especially chronic) or in a tissue,
 CC an allergic response in a subject, exposure of a subject to a pathogen
 CC or sterile inflammatory disease, by contacting a tissue having
 CC inflammation with an agent that modulates the expression of gene(s)
 CC from Gs in the tissue. M1 is useful for detecting GCA; M2 is useful for
 CC modulating Gc; M3 is useful for screening an agent capable of modulating
 CC GCA preferably in an inflammation in a tissue; M4 is useful for
 CC detecting an inflammation (especially chronic) in a tissue, an allergic
 CC response in a subject, exposure of a subject to a pathogen or sterile
 CC inflammatory disease (e.g. psoriasis, rheumatoid arthritis,
 CC glomerulonephritis, asthma, thrombosis, cardiac reperfusion injury, renal
 CC reperfusion injury, AIDS, adult respiratory distress syndrome,
 CC inflammatory bowel disease, Crohn's disease, ulcerative colitis,
 CC periodontal disease, also bacterial infection, viral infection,
 CC parasitic infection, protozoal infection, fungal infection and M5 is
 CC useful for treating one of the above conditions. The present
 CC sequence represents a gene differentially expressed in granulocytes.
 CC Note: the sequence data for this patent did not form part
 CC of the printed specification, but was obtained in electronic
 CC format directly from WIPO at
 CC http://wipo.int/pub/published_pct_sequences.

XX Sequence 3044 BP; 826 A; 687 C; 855 G; 675 T; 1 other.

Alignment Scores:

Prod. No.: 0.0057 Length: 3044
 Score: 65.00 Matches: 13
 Percent Similarity: 100.00% Conservative: 0
 Best Local Similarity: 100.00% Mismatches: 0
 Query Match: 100.00% Indels: 0
 DE: 24 Caps: 0

US-09-856-070-19 (1-13) x ARK#4562 (1-3044)

QY 1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
 |||||||
 IB 1147 AAGGAGGAGTTCATGCTGGCGCTGCCAGACATACAGGAG 1185
 |||||||

RESULT 6
 ABN97223
 ID ABN97223 standard; DNA: 3044 BP.
 XX
 AC ABN97223;
 XX
 DT 13-AUG-2002 (first entry)
 XX
 DE Gene #3721 used to diagnose liver cancer.
 XX
 KW Gene, liver cancer, ds, hepatocellular carcinoma; hepatotropic;
 KW metastatic liver tumor; cytostatic, expression profile; disease state;
 KW disease progression; drug toxicity; drug efficacy; drug metabolism.
 XX
 OS Homo sapiens.
 XX
 PN W0200229103-A2.
 XX
 PD 11-APR-2002.
 XX
 PF 02-OCT-2001; 2001WO-053589.
 XX
 PR 02-OCT-2000; 2000US-237054P.
 XX
 PA (GENE-) GENE LOGIC INC.
 XX
 PI Horne D, Alvares C, Peres-Da-Silva S, Vockley JG;
 XX W01: 2002 426119/45.
 XX
 XX Diagnosing and detecting the progression of liver cancer,
 PT hepatocellular carcinoma or metastatic liver tumor in a patient,
 PT involves detecting the level of expression of two or more genes in a
 PT liver tissue sample.

XX
 PS Claim 1; SEQ ID NO 3721; 298pp; English.

XX The invention relates to a novel method for diagnosing and detecting the
 CC progression of liver cancer, hepatocellular carcinoma or metastatic liver
 CC tumor in a patient, and differentiating metastatic liver cancer from
 CC hepatocellular carcinoma in a patient, involving detecting the level of
 CC expression of two or more genes represented in ARN93503-ARN97455 in a
 CC tissue sample. The method of the invention has hepatotropic, and
 CC cytostatic activity. The method is useful for diagnosing and detecting
 CC the progression of liver cancer, hepatocellular carcinoma and metastatic
 CC liver carcinoma in a patient. The method is useful for identifying
 CC expression profiles which serve as useful diagnostic markers as well as
 CC markers that can be used to monitor disease states, disease progression,
 CC drug toxicity, drug efficacy and drug metabolism.
 CC Note: The sequence data for this patent did not form part of the printed
 CC specification, but was obtained in electronic format directly from WIPO
 CC at http://wipo.int/pub/published_pct_sequences.

XX Sequence 3044 BP; 826 A; 687 C; 855 G; 675 T; 1 other;

Alignment Scores:

Prod. No.: 0.0057 Length: 3044
 Score: 65.00 Matches: 13
 Percent Similarity: 100.00% Conservative: 0

Best Local Similarity: 100.00% Mismatches: 0
 Query Match: 100.00% Indels: 0
 DB: 24 Gaps: 0
 US-09-856-070-19 (1-13) x ARK09723 (1-3044)

QY 1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
 DQ 1147 AAGGAGGAGTTCATGCTGGCGCTGACGAGCTATGAGGAG 1185

RESULT 7

ABK09792
 ID ARK09792 standard; cDNA: 3047 BP

XX AC ARK09792;

XX 14-MAR-2002 (first entry)

XX Human ovarian tumour protein encoding cDNA #325

XX Human; ovarian tumour protein, cancer, cytostatic, immunostimulant, ss.
 KW gene therapy; CD4+ T cell, CD8+ T cell, PCR primer.

XX Homo sapiens.

XX W0200190154-A2

XX 29-NOV-2001.

XX 23-MAY-2001; 2001WO/051680e

XX 24-MAY-2000; 2000US-207107p

XX 13-JUN-2000; 2000US-211457p

XX 21-JUN-2000; 2000US-211474p

XX 03-AUG-2000; 2000US-232288p

XX 01-MAR-2001; 2001US-0272790p

XX (CORI-) CORIXA CORP.

XX Xu J, Mitcham JL, Harlocker SL, Dillon DC, Socrist R, Lodes MJ;

PI Alidate PA, Fling SP, Mannion J, Benson DR, Carter D;

XX WPI: 2002-007641/13

XX New isolated polynucleotide encoding polypeptide comprising portion of

XX ovarian tumour protein, useful for detection, diagnosis and therapy of

XX human ovarian cancer

XX Claim 1; Page 269 270; 285pp; English.

XX The invention relates to an isolated polynucleotide encoding a
 CC polypeptide comprising a portion of an ovarian tumour protein. The
 CC sequences of the invention are useful for stimulating an immune response
 CC and for treating ovarian cancer in a patient. An antigen presenting cell
 CC that expresses the sequences is useful for treating ovarian cancer by
 CC incubating CD4+ and/or CD8+ T cells isolated from a patient. The T cells
 CC can then be proliferated and administered to the patient to inhibit the
 CC development of cancer. The DNA sequences are useful as probes or primers
 CC for nucleic acid hybridisation, to direct expression of a polypeptide in
 CC appropriate host cells. Detecting the presence of a cancer in a patient
 CC involves obtaining a biological sample from the patient, contacting the
 CC biological sample with an agent that binds to the protein, detecting the
 CC amount of protein that binds to the agent, comparing the amount of
 CC protein to a predetermined cut-off value and determining the presence of
 CC cancer. Sequences ARK09464-ARK09802 represent PCR primers and cDNA
 CC molecules encoding ovarian tumour proteins of the invention

XX Sequence 3047 BP; 828 A; 687 C; 856 G; 675 T; 1 other;

Alignment Scores: Alignment Scores: 3047
 Pred. No.: 0.0057 Length: 14
 Score: 65.00 Matches: 13
 Percent Similarity: 100.00% Conservative: 0

Best Local Similarity: 100.00% Mismatches: 0
 Query Match: 100.00% Indels: 0
 DB: 24 Gaps: 0
 US-09-856-070-19 (1-13) x ARK09792 (1-3047)

QY 1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
 DQ 1147 AAGGAGGAGTTCATGCTGGCGCTGACGAGCTATGAGGAG 1185

RESULT 8

ABK08182
 ID ARK08182 standard; cDNA: 3072 BP

XX AC ARK08182;

XX 18 SEP 2002 (first entry)

XX Human osteoblast differentiation related cDNA SEQ ID NO 89.

XX Human; osteoblast, stem cell differentiation, bone tissue deposition;
 KW osteoporosis, osteopathic; ss.
 XX Homo sapiens.

XX W0200250301-A2.

XX 27-JUN-2002.

XX 18 DEC 2001; 2001WO/3548276.

XX 18 DEC 2000; 2000US-251822P.

XX 24-APR-2001; 2001US-285691P.

XX (GENE-) GENE LOGIC INC.

XX (PROM) PROMETER & GAMBLE CO.

XX Ji D, Axelrod DW, Cook JS, Jaiswal N, Einstein R, Houghlon A;

PI Mertz L;

XX WPI: 2002-557663/59.

XX The invention relates to their expression profiles associated with osteoblast
 PT differentiation for screening modulators bone formation, for diagnosing
 PT or treating e.g. osteoporosis, or as markers for the differentiation
 PT process

XX Claim 1; SEQ ID NO 89; 78pp + Sequence Listing; English.

XX The invention relates to genes and their expression profiles are used

XX for:
 CC (a) screening modulators of precursor stem cell differentiation into
 CC osteoblasts, or bone tissue deposition;

XX (b) diagnosing abnormal deposition of bone tissue, abnormal rate of
 CC osteoblast formation or osteoporosis; or

XX (c) treating or monitoring treatment of the conditions cited in (b), or
 CC monitoring the progression of bone tissue deposition.

XX Specific conditions include postmenopausal osteoporosis, glucocorticoid
 CC osteoporosis or male osteoporosis, osteopenia, osteodystrophy,
 CC drug-induced abnormalities in bone formation or bone loss, conditions
 CC that involve altered bone metabolism (e.g. idiopathic juvenile
 CC osteoporosis), skeletal disease linked to breast cancer, mastocytosis,
 CC Paget's syndrome of fibrous dysplasia, the present sequence is that of an
 CC osteoblast differentiation associated cDNA marker of the invention.

CC Note: The sequence data for this patent did not form part of the printed
 CC specification, but was obtained in electronic format directly from WIPO
 CC at ftp.wipo.int/pub/published_pat_sequences.

XX Sequence 3072 BP; 846 A; 688 C; 868 G; 670 T; 0 other;

Alignment Scores: Alignment Scores: 3072
 Pred. No.: 0.0057 Length: 13
 Score: 65.00 Matches: 13

Percent Similarity: 100.00%
Best Local Similarity: 100.00%
Query Match: 24
DH: 0

Score: 65.00
Percent Similarity: 100.00%
Best Local Similarity: 100.00%
Query Match: 21
DH: 0

US-09-856-070-19 (1-13) x AAG88182 (1-3072)

QY 1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
|||||
DB 1163 AAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 1201

RESULT 9

ID AAG98114 standard; cDNA: 3115 BP.

AC AAG98114;

DT 09-MAR-2001 (first entry)

Human colon cancer antigen nucleotide sequence SEQ ID NO:123.

Human; colon cancer; colon cancer antigen; diagnosis; detection;
identification; cytostatic; cardioactive; neuroprotective; vulnary;
immunomodulatory; muscular; anaerobic; gastrointestinal;
neurotropic; antineoplastic; antibacterial; gene therapy; wound;
neural disorder; immune system disorder; muscular disorder;
reproductive disorder; gastrointestinal disorder; renal disorder;
infectious disease; cardiovascular disorder; ss.

OS Homo sapiens.

XX W0200055451 A1.

PD 21-SEP-2000.

PF 08-MAR-2000; 2000W-US05883

XX 12-MAR-1999; 9905-0124270.

XX (HUMA-) HUMAN GENOME SCI INC.

XX Rosen CA, Ruben SM;

XX WPI: 2000 587544/55

XX P-PSDB; AAB53356.

Colon cancer associated gene sequences, referred to as colon cancer
antigens, useful for the treatment, prevention, and diagnosis of colon
disorders such as colon cancer

Claim 1: Page 554-560; 2104pp; English

AAC97991 to AAC98763 encodes the human colon cancer associated proteins
called human colon cancer antigens, given in AAC98324 to AAC98406. The
human colon cancer antigen can have cytostatic, cardioactive, muscular;
neuroprotective immunomodulatory anaerobic; gastrointestinal, muscular;
vulnary, nephrotropic, antineoplastic and antibacterial activities, and
can be used in gene therapy. The colon cancer antigen polypeptides,
proteins and antibodies to the proteins are useful for the prevention,
treatment and diagnosis of colon disorders, such as colon cancer. The
polynucleotides may be used in diagnostics and research, such as for
chromosome identification, and as hybridisation probes. The proteins
may also be used to prevent diseases such as neural disorders, immu-
ne system disorders, muscular disorders, reproductive disorders,
gastrointestinal disorders, wounds, renal disorders, infectious
diseases, and cardiovascular disorders. AAC98764 to AAC98772 and
AAB54007 represent sequences used in the exemplification of the present
invention.

Sequence 3115, 400, 873 A, 666 C, 872 G, 670 T, 4 other;

Alignment Scores: 0.00585 Length: 3115

Pred. No.:

US-09-856-070-19 (1-13) x AAC98113 (1-3115)
QY 1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
|||||
DB 1179 AAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 1217

RESULT 10

AAK70537/C

ID AAK70537 standard; DNA: 11445 BP.

XX AAK70537;

AC AAK70537;

DT 06-NOV-2001 (first entry)

Human immune/haematopoietic antigen genomic sequence SEQ ID NO:25349.
Human; immune; haematopoietic; immune/haematopoietic antigen; cancer;
cytostatic; gene therapy; vaccine; metastasis; ds.
Homo sapiens.
OS
XX W0200157182 A2.

PD 09-AUG-2001.

XX 17-JAN-2001; 2001W-US01354.

XX 31-JAN-2000; 2000US-0174065.

XX 04-FEB-2000; 2000US-0180628.

XX 24-FEB-2000; 2000US-0184664.

XX 02-MAR-2000; 2000US-0186350.

XX 16-MAR-2000; 2000US-0189874.

XX 17-MAR-2000; 2000US-0190076.

XX 18-APR-2000; 2000US-0198123.

XX 19-MAY-2000; 2000US-0205515.

XX 07-JUN-2000; 2000US-0209467.

XX 28-JUN-2000; 2000US-0214886.

XX 30-JUN-2000; 2000US-0215145.

XX 07-JUL-2000; 2000US-0216647.

XX 07-JUL-2000; 2000US-0216840.

XX 11-JUL-2000; 2000US-0217487.

XX 14-JUL-2000; 2000US-0218290.

XX 28-JUL-2000; 2000US-0220963.

XX 14-AUG-2000; 2000US-0220964.

XX 14-AUG-2000; 2000US-0224518.

XX 14-AUG-2000; 2000US-0224519.

XX 14-AUG-2000; 2000US-0225213.

XX 14-AUG-2000; 2000US-0225214.

XX 14-AUG-2000; 2000US-0225266.

XX 14-AUG-2000; 2000US-0225267.

XX 14-AUG-2000; 2000US-0225268.

XX 14-AUG-2000; 2000US-0225270.

XX 14-AUG-2000; 2000US-0225447.

XX 14-AUG-2000; 2000US-0225757.

XX 14-AUG-2000; 2000US-0225758.

XX 14-AUG-2000; 2000US-0225759.

XX 18-AUG-2000; 2000US-0226279.

XX 22-AUG-2000; 2000US-0226881.

XX 22-AUG-2000; 2000US-0226868.

XX 22-AUG-2000; 2000US-0227182.

XX 23-AUG-2000; 2000US-0227009.

XX 30-AUG-2000; 2000US-0228324.

XX 01-SEP-2000; 2000US-0229287.

XX 01-SEP-2000; 2000US-0229343.

XX 01-SEP-2000; 2000US-0229344.

XX 01-SEP-2000; 2000US-0229345.

XX AAS93352;
 XX 14-FEB-2002 (first entry)
 XX DNA encoding novel human diagnostic protein #29156.
 XX Human; chromosome mapping; gene mapping; gene therapy; forensic;
 XX food supplement; medical imaging; diagnostic; genetic disorder; ss.
 XX Homo sapiens.
 XX W0200175067-A2.
 XX 11-OCT-2001.
 XX 40-MAR-2001; 2001WO-0508631.
 XX 41 MAR-2000; 2000US-0540217.
 XX 24-AUG-2000; 2000US 0649167.
 XX (HUSE-) HYSRQ INC.
 XX Demanac RT, Liu C, Tang YT;
 XX WPI: 2001 649362/74.
 XX P-PSDB; ARG29165.
 XX New isolated polynucleotide and encoded polypeptides, useful in
 XX diagnostics, forensics, gene mapping, identification of mutations
 XX responsible for genetic disorders or other traits and to assess
 XX biodiversity.
 XX Claim 1: SEQ ID NO 29156; 103pp; English.
 XX The invention relates to isolated polynucleotide (I) and
 XX polypeptide (II) sequences. (I) is useful as hybridisation probes,
 XX polymerase chain reaction (PCR) primers, oligomers, and for chromosome
 XX and gene mapping, and in recombinant production of (II). The
 XX polynucleotides are also used in diagnostics as expressed sequence tags
 XX for identifying expressed genes. (I) is useful in gene therapy techniques
 XX to restore normal activity of (II) or to treat disease states involving
 XX (II). (II) is useful for generating antibodies against it, detecting or
 XX quantitating a polypeptide in tissue, as molecular weight markers and as
 XX a food supplement. (II) and its binding partners are useful in medical
 XX imaging of sites expressing (II). (I) and (II) are useful for treating
 XX disorders involving aberrant protein expression or biological activity.
 XX The polypeptide and polynucleotide sequences have applications in
 XX diagnostics, forensics, gene mapping, identification of mutations
 XX responsible for genetic disorders or other traits to assess biodiversity
 XX and to produce other types of data and products dependent on DNA and
 XX amino acid sequences. AAS64197 AAS94364 represent novel human
 XX diagnostic coding sequences of the invention.
 XX Note: The sequence data for this patent did not appear in the printed
 XX specification, but was obtained in electronic format directly from WIPO
 XX at ftp.wipo.int/pub/published_pct_sequences.

XX Sequence 1447 BP; 347 A; 356 C; 383 G; 361 T; 0 other;

Alignment Scores:	139	Length:	1447
Pred. No.:	41.00	Matches:	8
Score:	90.91%	Conservative:	2
Percent Similarity:	72.73%	Mismatches:	0
Best Local Similarity:	63.08%	Indels:	0
Query Match:	24	Gaps:	0

US-09 856-070 19 (1-13) x AAS93352 (1-1447)

QY 2 GluGluGluGluMetLeuArqLeuGlnAspTyrGluGln 12

Ub 979 AAGAGTTGATGTCAGGAGAGAGATTACGAG 1011

RESULT 12
 ABQ88140/C
 ID ABQ88140 standard; cDNA; 66494 BP.
 XX AC ABQ88140;
 XX 18-SEP-2002 (first entry)
 XX Human osteoblast differentiation related cDNA SEQ ID NO 47.
 XX Human; osteoblast; stem cell differentiation; bone tissue deposition;
 XX osteoporosis; osteopathic; ss.
 XX Homo sapiens.
 XX W0200205040 A2.
 XX 27-JUN-2002.
 XX 18-DEC-2001; 2001WO-0548476.
 XX 18-DEC-2000; 2000US 255882P.
 XX 24-APR-2001; 2001US 285691P.
 XX (GENE-) GENE LOGIC INC.
 XX (PROC) PROCTER & GAMBLE CO.
 XX JI D, Axelrod DW, Cook JS, Jaiswal N, Einstein R, Houghton A;
 XX Mortz L;
 XX WPI: 2002-557663/59.
 XX Use of genes and their expression profiles associated with osteoblast
 XX differentiation for screening modulators bone formation, for diagnosing
 XX or treating e.g. osteoporosis, or as markers for the differentiation
 XX process.
 XX Claim 1: SEQ ID NO 47; 78pp + Sequence Listing; English.
 XX The invention relates to genes and their expression profiles are used
 XX for:
 XX (a) screening modulators of precursor stem cell differentiation into
 XX osteoblasts, or bone tissue deposition;
 XX (b) diagnosing abnormal deposition of bone tissue, abnormal rate of
 XX osteoblast formation or osteoporosis; or
 XX (c) treating or monitoring treatment of the conditions cited in (b), or
 XX monitoring the progression of bone tissue deposition.
 XX Specific conditions include postmenopausal osteoporosis, glucocorticoid
 XX osteoporosis or male osteoporosis, osteopenia, osteodystrophy,
 XX drug-induced abnormalities in bone formation or bone loss, conditions
 XX that involve altered bone metabolism (e.g. idiopathic juvenile
 XX osteoporosis), skeletal disease linked to breast cancer, mastocytosis,
 XX Fanconi syndrome or fibrous dysplasia. The present sequence is that of an
 XX osteoblast differentiation associated cDNA marker of the invention.
 XX Note: The sequence data for this patent did not form part of the printed
 XX specification, but was obtained in electronic format directly from WIPO
 XX at ftp.wipo.int/pub/published_pct_sequences.

XX Sequence 66494 BP; 17680 A; 14908 C; 14466 G; 20440 T; 0 other;

Alignment Scores:	1248+04	Length:	66494
Pred. No.: <td>41.00 <td>Matches: <td>8</td> </td></td>	41.00 <td>Matches: <td>8</td> </td>	Matches: <td>8</td>	8
Score: <td>84.62%</td> <td>Conservative: <td>4</td> </td>	84.62%	Conservative: <td>4</td>	4
Percent Similarity: <td>61.54%</td> <td>Mismatches: <td>2</td> </td>	61.54%	Mismatches: <td>2</td>	2
Best Local Similarity: <td>63.08%</td> <td>Indels: <td>0</td> </td>	63.08%	Indels: <td>0</td>	0
Query Match: <td>24</td> <td>Gaps: <td>0</td> </td>	24	Gaps: <td>0</td>	0

US-09-856-070-19 (1-13) x ABQ88140 (1-66494)

QY 1 LysGluGluGluMetLeuArqLeuGlnAspTyrGluGln 14

Db 57190 AAGAGTTCAGGAGAGAGATTACGAG 57152


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RESULT 13
ABA74089/c
ID ABA74089 standard; DNA: 224 BP.
XX
AC ABA74089;
XX
DI 01-FEB-2002 (first entry)
XX
DE Human foetal liver single exon nucleic acid probe #22304
XX
KW Human; foetal liver; gene expression, single exon nucleic acid probe; ss.
XX
OS Homo sapiens.
XX
PN WO200157277-A2.
XX
PO 09-AUG-2001.
XX
PF 30-JAN-2001; 2001WO-US00669.
XX
PR 04-FEB-2000; 2000US-0180312.
PR 26-MAY-2000; 2000US-0207456.
PR 30-JUN-2000; 2000US-0608408.
PR 03-AUG-2000; 2000US-0612466.
PR 21-SEP-2000; 2000US-0234687.
PR 27-SEP-2000; 2000US-0236359.
PR 04-OCT-2000; 2000GB-0024263.
XX
PA (MOL.) MOLECULAR DYNAMICS INC.
XX
PI Penn SG, Hanzel DK, Chen W, Rank DR;
XX
WI: 2001-483447/52.
XX
PT Human genome-derived single exon nucleic acid probes useful for
    analyzing gene expression in human foetal liver.
XX
PS Claim 4; SEQ ID NO 22304; 60pp + sequence listing, English.
XX
CC The invention relates to a single exon nucleic acid probe for
    measuring human gene expression in a sample derived from human foetal
    liver. The single exon nucleic acid probes may be used for predicting,
    measuring and displaying gene expression in samples derived from human
    foetal liver. The present sequence is a single exon nucleic acid
    probe of the invention.
CC Note: The sequence data for this patent did not form part of the
    printed specification, but was obtained in electronic format directly
    from WIPO at ftp.wipo.int/pub/published_pat_sequences
XX
SQ Sequence 224 BP; 58 A; 63 C; 45 G; 58 T; 0 other.

Alignment Scores:
Pred. No.: 24.6 Length: 224
Score: 40.00 Matches: 7
Percent Similarity: 84.62% Conservative: 4
Best Local Similarity: 53.85% Mismatches: 2
Query Match: 61.54% Indels: 0
Gaps: 0

US-09-856-070-19 (1-13) x ABA74089 (1-224)
QY 1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
||||| ||||| ||||| |||||
Db 57 AAGAGAAATCTGCTGCTGGAACTCGGTAAATATAGAGAA 19

RESULT 14
AAK22545/c
ID AAK22545 standard; DNA: 224 BP.
XX
AC AAK22545;
XX
DI 05-NOV-2001 (first entry)
XX
DE Human foetal liver single exon nucleic acid probe #9897.
XX
KW Human; foetal liver; gene expression, single exon nucleic acid probe; ss.
XX
OS Homo sapiens.
XX
PN WO200157277-A2.
XX
PO 09-AUG-2001.
XX
PF 30-JAN-2001; 2001WO-US00667.
XX
PR 04-FEB-2000; 2000US-0180312.
PR 26-MAY-2000; 2000US-0207456.
PR 30-JUN-2000; 2000US-0608408.
PR 03-AUG-2000; 2000US-0612466.
PR 21-SEP-2000; 2000US-0234687.
PR 27-SEP-2000; 2000US-0236359.
PR 04-OCT-2000; 2000GB-0024263.
XX
PA (MOL.) MOLECULAR DYNAMICS INC.
XX
PI Penn SG, Hanzel DK, Chen W, Rank DR;
XX
WI: 2001-483447/52.
XX
PT Single exon nucleic acid probes for analyzing gene expression in human
    brains.
XX
PS Example 4; SEQ ID NO 22304; 60pp + sequence listing, English.
XX
CC The present invention provides a number of single exon nucleic acid
    probes which are derived from genomic sequences expressed in the human
    brain. They can be used to measure gene expression in brain cell samples,
    which may enable the diagnosis and improved treatment of nervous system
    diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia,
    epilepsy and cancers. The present sequence is one of the probes of the
    invention.
XX
SQ Sequence 224 BP; 58 A; 63 C; 45 G; 58 T; 0 other.

Alignment Scores:
Pred. No.: 24.6 Length: 224
Score: 40.00 Matches: 7
Percent Similarity: 84.62% Conservative: 4
Best Local Similarity: 53.85% Mismatches: 2
Query Match: 61.54% Indels: 0
Gaps: 0

US-09-856-070-19 (1-13) x AAK22545 (1-224)
QY 1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
||||| ||||| ||||| |||||
Db 57 AAGAGAAATCTGCTGCTGGAACTCGGTAAATATAGAGAA 19

RESULT 15
AAH61592/c
ID AAH61592 standard; DNA: 520 BP.
XX
AC AAH61592;
XX
DI 01-FEB-2002 (first entry)
XX
DE Human foetal liver single exon nucleic acid probe #9897.
XX
KW Human; foetal liver; gene expression, single exon nucleic acid probe; ss.
XX
OS Homo sapiens.
XX
PN WO200157277-A2.

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XX 09 AUG-2001.
XX 40 JAN-2001: 2001RG-US000669.
XX 04 FEB-2000: 2000US-0180312.
XX 26 MAY-2000: 2000US-0207456.
XX 40 JUN-2000: 2000US-0608408.
XX 03 AUG-2000: 2000US-0632366.
XX 21 SEP-2000: 2000US-0244687.
XX 27 SEP-2000: 2000US-0236459.
XX 04 OCT-2000: 2000US-0024263.
XX (MOLE) MOLECULAR DYNAMICS INC.
XX Penn St, Hanzel DK, Chen W, Rank DR;
XX WPI: 2001-48447/52.
XX Human genome-derived single exon nucleic acid probes useful for
XX analyzing gene expression in human fetal liver.
XX Claim 1: SEQ ID NO 9897; 639pp + sequence listing; English.
XX the invention relates to a single exon nucleic acid probe for
XX measuring human gene expression in a sample derived from human fetal
XX liver. The single exon nucleic acid probes may be used for predicting,
XX measuring and displaying gene expression in samples derived from human
XX fetal liver. The present sequence is a single exon nucleic acid
XX probe of the invention.
XX Note: The sequence data for this patent did not form part of the
XX printed specification, but was obtained in electronic format directly
XX from WPI at ftp.wipo.int/pub/published_pat_sequences.
XX Sequence 520 BP; 135 A; 145 C; 88 G; 152 T; 0 other;

Alignment Scores:
Pred. No.: 66.1 Length: 520
Score: 40.00 Matches: 7
Percent Similarity: 84.62% Conservative: 4
Best Local Similarity: 53.85% Mismatches: 2
Query Match: 61.54% Indels: 0
Gaps: 0

US 09 856-070 19 (1-13) x ABA51592 (1-520)
QY 1 LysGluLeuLeuMetLeuArgLeuGluAspTyrGluGlu 13
DB 150 AAGGAGAAATCTGTTCCTCGAATCTCCGTAACATGAGGAA 112

Search completed: January 16, 2003, 17:19:45
Job time : 218.954 secs

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